

Figure 1a (2)

Figure 1a (2)

[illegible]

Figure 1a (3)

Cslseq	(2048)	agaatttgtaagttctctgaagatggcaaaagataacgataaatttcgcattaaagcgctacatctccaataacactatcaaccccgctttgacccacgcggtgagcgagtatatcgcgctc
Kukkaseq	(1890)tgc.....t.c.c.....t.c.a.....t.....c.t.t.....c.a.....t.g.a.....t.....
Ds4seq	(1844)a.tgc.....t.c.c.....t.c.a.....t.....c.t.....c.t.....t.....
2301SEQ	(1843)tgc.....t.c.c.....c.a.....t.....c.t.....c.t.....t.....
390seq	(1845)tgc.....t.c.c.....c.a.....t.....c.....gga.cg.g.t.t.ga.ttct.c.g.g.....
H. felis com	(1920)	g.....gcgct.gaaa.g.aaagggc.....c.c.....c.a.....t.g.t.....c.....a.ga.cg.t.t.ga.t.....g.a.t.....
H. pylori co	(4533)cgct.gaaa.....aaagggc.....c.c.....a.g.c.a.....t.g.t.....c.....t.ca.c.a.....a.ttct.a.g.g.....
H. heilmanni	(2081)cgct.g.....aaagggc.c.....c.c.....c.a.....t.t.....c.....t.ca.c.a.....a.ttct.a.g.g.....
Cslseq	(2168)	tgtggaagaggccaagatccgcgactgtgtgtggtgggaatcctgccttttttggcgtaaaaccccaaaatcgtgatcaaaagcggtatgggtgtcttcttgaatggcggtattcttaagcgc
Kukkaseq	(2010)a.....t.t.....t.c.....t.g.....t.g.....g.....c.c.....
Ds4seq	(1964)a.....t.t.....t.c.....t.g.....t.g.....t.....c.....c.c.....
2301SEQ	(1963)t.....t.c.....t.g.....t.g.....t.....c.....t.....g.c.t.....
390seq	(1965)t.....ata.....c.c.....c.t.....g.g.t.c.....a.t.g.....t.ga.t.t.g.....at.ta.t.cgc.....c.....g.c.t.....
H. felis com	(2040)t.....t.....c.c.....c.t.....g.g.t.c.....a.t.g.....t.ga.t.t.g.....at.ta.t.cgc.....aagcc.....g.g.....
H. pylori co	(4653)	a.a.....t.....ag.g.t.....at.....g.a.a.c.....g.....c.g.a.c.....t.ca.t.cgc.t.....c.....g.c.....
H. heilmanni	(2201)a.....t.t.ata.....t.....c.t.....gc.....g.c.....a.t.....c.g.a.c.....t.....tccaagagtcgcctatgaaaatggcggtgaaaga
Cslseq	(2288)	gtctgtgccactccccaacggtttattaccgcggaatgtttgggcataccggaaggcgaaatttgacaccagcatcacttttgttccaaagtcgcctatgaaaatggcggtgaaaga
Kukkaseq	(2130)t.....t.g.....t.g.....t.....t.....t.....
Ds4seq	(2084)t.a.t.g.....t.g.....t.....t.....t.....t.....
2301SEQ	(2083)a.t.g.....t.g.....c.....t.g.aaac.....c.....at.....c.g.c.....cg.t.ca.ggca.ga.c.....
390seq	(2085)c.....g.t.g.....t.....a.c.t.g.aaac.....c.....at.....c.g.c.....cg.t.ca.ggca.ga.c.....
H. felis com	(2160)a.t.....c.t.g.c.c.....t.....c.ct.....t.t.a.t.....ac.tg.a.a.....g.tc.cg.t.....c.a.a.t.....
H. pylori co	(4773)	t.....a.c.t.c.a.....a.a.....c.ct.....t.t.a.t.....ac.tg.a.a.....g.tc.cg.t.....c.a.a.t.....
H. heilmanni	(2321)	t.....a.c.....c.g.c.....c.c.c.....a.c.....a.c.....c.g.t.c.....g.t.....c.a.t.....c.c.....
Cslseq	(2408)	aaagtggttagcgccaagtctaccgggtcaaaaactgcgttaacatcaccagaagaagactcaagttcaacgacaaaacgcaaaatcacccgtcgatccgaaaaccttcgaggt
Kukkaseq	(2250)a.a.t.....a.g.g.c.c.g.....g.....t.....t.....t.....t.....t.....
Ds4seq	(2204)a.a.....a.g.g.....c.g.....c.....t.....a.....a.....g.....gc.t.....t.....t.....
2301SEQ	(2203)a.a.....a.g.g.....c.g.....c.....t.....a.....a.....g.....gc.t.....t.....t.....
390seq	(2170)a.....a.g.g.....c.g.....c.....t.....t.a.g.....c.....a.....tggtg.c.c.t.tgat.a.c.tg.....ata.....
H. felis com	(2280)	g.a.a.gc.....t.gcg.ca.cg.a.g.....t.c.t.....t.a.g.....c.....a.....tggtg.c.c.t.tgat.a.c.tg.....ata.....
H. pylori co	(4893)	g.at.a.ac.t.aa.....gt.g.....t.a.a.t.....t.a.....a.gc.a.....ct.t.tc.c.tgaa.a.a.tg.....t.a.c.t.....
H. heilmanni	(2441)	cg.t.....gc.aa.agtt.gt.g.t.g.....c.....a.....tgct.c.c.c.gaa.a.c.tg.....a.a.a.....
Cslseq	(2528)	ctttgtagatggcaaacctctgcaccttaaacccactcgcaagtcgctctagccccagcgctacacttttctctag
Kukkaseq	(2370)g.....tg.....a.....t.....t.....
Ds4seq	(2324)g.....tg.....a.....t.....t.....
2301SEQ	(2323)g.....tg.....a.....t.....t.....
390seq	(2170)c.t.....a.....t.....t.....
H. felis com	(2400)	gaaa.g.....gaggtg.....g.ag.agatg.t.agc.....g.a.tt.t.a.g.....
H. pylori co	(5013)	g.c.g.....gaagta.t.....ag.aaata.....agct.g.g.a.t.tt.gca.t.....
H. heilmanni	(2561)	gaaa.g.....cgaagtt.....cc.tg.gg.tgaca.t.agc.....a.a.t.....t.acc.a.....

Figure 1a (4)

Figure 1b

ALIGNED SEQUENCES

Figure 1c (1)

Figure 1c (1)

THE T40 "THE T40"

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ureYCS1      ( 480) hhgkakfdtsitfsvkvayengvkeklglergvlpvknrcrnitkkafkfnctakitvdpktfevfdgklctskptsqvplaqrytff
ureYkuka      ( 480) .....k.....n.....h.....e.....
ureYDS4      ( 480) .....k.....k.....h.....a.e.....
ureY2301      ( 480) .....k.....k.....h.....a.e.....rvss
ureY390      ( 480) .....n.....qa..ka.i..e..d.aap.....l....v..h.d.n.e.yk.k....ev...aadelS...l.nl.
B felis      ( 481) .....Y.an.....qa..dk.i..e.....mq...t..h.e.n.e.yh.....ev...ank.S...lfsi.
B pylori      ( 481) .....n.....q.....i.he....q.v.....l....v..h.e.n.e.yk.k...nev..haadkls...l.nl.
B heilman      ( 480) .....n.....q.....i.he....q.v.....l....v..h.e.n.e.yk.k...nev..haadkls...l.nl.

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Figure 1c (2)

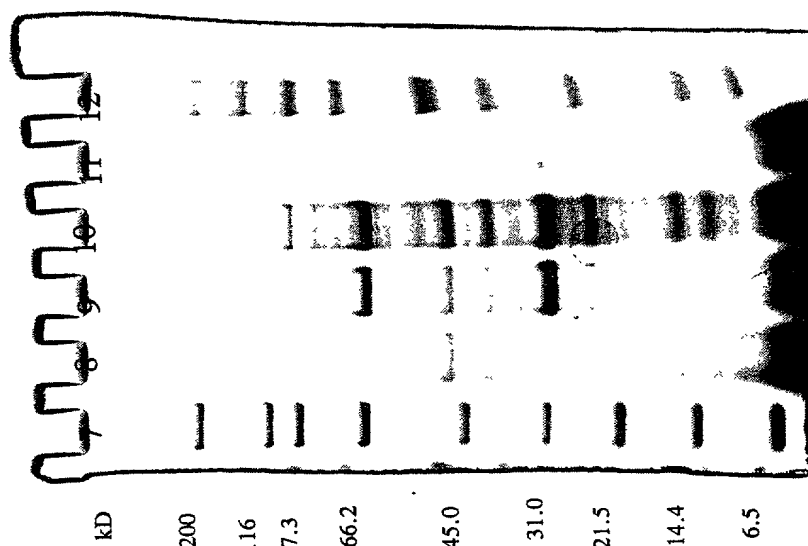


Figure 2